

SEQUENCE LISTING

Chen, Jin-Long Tian, Hui Zhong, Wendy Wen Tularik Inc. <120> Receptor Ligands and Methods of Modulating Receptors <130> 018781-009530US <140> US 10/719,692 <141> 2003-11-21 <150> US 60/421,142 <151> 2002-11-25 <150> US 60/444,153 <151> 2003-01-30 <160> 9 <170> PatentIn Ver. 2.1 <210> 1 <211> 1164 <212> DNA <213> Homo sapiens <223> human G-protein coupled receptor TGR4a (HM74) <400> 1 atgaatcggc accatctgca ggatcacttt ctggaaatag acaagaagaa ctgctgtgtg 60 ttccgagatg acttcattgc caaggtgttg ccgccggtgt tgggggctgga gtttatcttt 120 gggcttctgg gcaatggcct tgccctgtgg attttctgtt tccacctcaa gtcctggaaa 180 tccagccgga ttttcctgtt caacctggca gtagctgact ttctactgat catctgcctg 240 ccqttcqtqa tqqactacta tqtqcqqcqt tcaqactgqa actttqgqqa catcccttqc 300 cqqctqqtqc tcttcatqtt tgccatgaac cgccagggca gcatcatctt cctcacggtg 360 gtggcggtag acaggtattt ccgggtggtc catccccacc acgccctgaa caagatctcc 420 aattggacag cagccatcat ctcttgcctt ctgtggggca tcactgttgg cctaacagtc 480 cacctcctga agaagaagtt gctgatccag aatggccctg caaatgtgtg catcagcttc 540 agcatctgcc ataccttccg gtggcacgaa gctatgttcc tcctggagtt cctcctgccc 600 ctgggcatca tcctgttctg ctcagccaga attatctgga gcctgcggca gagacaaatg 660 gaccggcatg ccaagatcaa gagagccatc accttcatca tggtggtggc catcgtcttt 720 gtcatctgct tccttcccag cgtggttgtg cggatccgca tcttctggct cctgcacact 780 tcgggcacgc agaattgtga agtgtaccgc tcggtggacc tggcgttctt tatcactctc 840 agetteacet acatgaacag catgetggac ecegtggtgt actaettete cageccatee 900 tttcccaact tcttctccac tttgatcaac cgctgcctcc agaggaagat gacaggtgag 960 ccagataata accgcagcac gagcgtcgag ctcacagggg accccaacaa aaccagaggc 1020 gctccagagg cgttaatggc caactccggt gagccatgga gcccctctta tctgggccca 1080 acctcaaata accattccaa gaagggacat tgtcaccaag aaccagcatc tctggagaaa 1140 1164

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<211> 387

<212> PRT

<213> Homo sapiens

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Asp Leu Ala Phe Phe Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met

Leu Asp Pro Val Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Asn Phe

295

290

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Pro Asp Asn Asn Arg Ser Thr Ser Val Glu Leu Thr Gly Asp Pro Asn
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Lys Thr Arg Gly Ala Pro Glu Ala Leu Met Ala Asn Ser Gly Glu Pro
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Trp Ser Pro Ser Tyr Leu Gly Pro Thr Ser Asn Asn His Ser Lys Lys
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Gly His Cys His Gln Glu Pro Ala Ser Leu Glu Lys Gln Leu Gly Cys
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Cys Ile Glu
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Lys Thr Arg Gly Ala Pro Glu Ala Leu Met Ala Asn Ser Gly Glu Pro 345 Trp Ser Pro Ser Tyr Leu Gly Pro Thr Ser Pro 360 <210> 5 <211> 1041 <212> DNA <213> Homo sapiens <220> <223> human G-protein coupled receptor TGR183 <400> 5 atgtacaacg ggtcgtgctg ccgcatcgag ggggacacca tctcccaggt gatgccgccg 60 ctgctcattg tggcctttgt gctgggcgca ctaggcaatg gggtcgccct gtgtggtttc 120 tgcttccaca tgaagacctg gaagcccagc actgtttacc ttttcaattt ggccgtggct 180 gatttcctcc ttatgatctg cctgcctttt cggacagact attacctcag acgtagacac 240 tgggcttttg gggacattcc ctgccgagtg gggctcttca cgttggccat gaacagggcc 300 gggagcatcg tgttccttac ggtggtggct gcggacaggt atttcaaagt ggtccacccc 360 caccacgcgg tgaacactat ctccacccgg gtggcggctg gcatcgtctg caccctgtgg 420 gccctggtca tcctgggaac agtgtatctt ttgctggaga accatctctg cgtgcaagag 480 acggccgtct cctgtgagag cttcatcatg gagtcggcca atggctggca tgacatcatg 540 ttccagctgg agttctttat gcccctcggc atcatcttat tttgctcctt caagattgtt 600 tggagcctga ggcggaggca gcagctggcc agacaggctc ggatgaagaa ggcgacccgg 660 ttcatcatgg tggtggcaat tgtgttcatc acatgctacc tgcccagcgt gtctgctaga 720 ctctatttcc tctggacggt gcctcgagt gcctgcgatc cctctgtcca tggggccctg 780 cacataaccc tcaqcttcac ctacatgaac agcatgctgg atcccctggt gtattatttt 840 tcaagcccct cctttcccaa attctacaac aagctcaaaa tctgcagtct gaaacccaag 900 cagccaggac actcaaaaac acaaaggccg gaagagatgc caatttcgaa cctcggtcgc 960 aggagttgca tcagtgtggc aaatagtttc caaagccagt ctgatgggca atgggatccc 1020 cacattgttg agtggcactg a <210> 6 <211> 346 <212> PRT <213> Homo sapiens <220> <223> human G-protein coupled receptor TGR183 Met Tyr Asn Gly Ser Cys Cys Arg Ile Glu Gly Asp Thr Ile Ser Gln Val Met Pro Pro Leu Leu Ile Val Ala Phe Val Leu Gly Ala Leu Gly 20 Asn Gly Val Ala Leu Cys Gly Phe Cys Phe His Met Lys Thr Trp Lys Pro Ser Thr Val Tyr Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu 55 Met Ile Cys Leu Pro Phe Arg Thr Asp Tyr Tyr Leu Arg Arg Arg His

75

70

65

Trp Ala Phe Gly Asp Ile Pro Cys Arg Val Gly Leu Phe Thr Leu Ala 90 Met Asn Arg Ala Gly Ser Ile Val Phe Leu Thr Val Val Ala Ala Asp 105 Arg Tyr Phe Lys Val Val His Pro His His Ala Val Asn Thr Ile Ser Thr Arg Val Ala Ala Gly Ile Val Cys Thr Leu Trp Ala Leu Val Ile 135 Leu Gly Thr Val Tyr Leu Leu Leu Glu Asn His Leu Cys Val Gln Glu Thr Ala Val Ser Cys Glu Ser Phe Ile Met Glu Ser Ala Asn Gly Trp 170 His Asp Ile Met Phe Gln Leu Glu Phe Phe Met Pro Leu Gly Ile Ile Leu Phe Cys Ser Phe Lys Ile Val Trp Ser Leu Arg Arg Gln Gln 200 Leu Ala Arg Gln Ala Arg Met Lys Lys Ala Thr Arg Phe Ile Met Val Val Ala Ile Val Phe Ile Thr Cys Tyr Leu Pro Ser Val Ser Ala Arg 235 Leu Tyr Phe Leu Trp Thr Val Pro Ser Ser Ala Cys Asp Pro Ser Val 250 His Gly Ala Leu His Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met . 265 Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Lys Phe 275 Tyr Asn Lys Leu Lys Ile Cys Ser Leu Lys Pro Lys Gln Pro Gly His Ser Lys Thr Gln Arg Pro Glu Glu Met Pro Ile Ser Asn Leu Gly Arg 310 Arg Ser Cys Ile Ser Val Ala Asn Ser Phe Gln Ser Gln Ser Asp Gly

<210> 7

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<212> DNA

<213> Homo sapiens

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340

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His Leu Leu Lys Lys Lys Leu Leu Ile Gln Asn Gly Pro Ala Asn Val 165 170 175

Cys Ile Ser Phe Ser Ile Cys His Thr Phe Arg Trp His Glu Ala Met 180 185 190

Phe Leu Leu Glu Phe Leu Leu Pro Leu Gly Ile Ile Leu Phe Cys Ser 195 200 205

Ala Arg Ile Ile Trp Ser Leu Arg Gln Arg Gln Met Asp Arg His Ala 210 215 220

Lys Ile Lys Arg Ala Ile Thr Phe Ile Met Val Val Ala Ile Val Phe 225 230 235 240

Val Ile Cys Phe Leu Pro Ser Val Val Val Arg Ile Arg Ile Phe Trp 245 250 255

Leu Leu His Thr Ser Gly Thr Gln Asn Cys Glu Val Tyr Arg Ser Val
260 265 270

Asp Leu Ala Phe Phe Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met 275 280 285

Leu Asp Pro Val Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Asn Phe 290 295 300

Phe Ser Thr Leu Ile Asn Arg Cys Leu Gln Arg Lys Met Thr Gly Glu 305 310 315 320

Pro Asp Asn Asn Arg Ser Thr Ser Val Glu Leu Thr Gly Asp Pro Asn 325 330 335

Lys Thr Arg Gly Ala Pro Glu Ala Leu Met Ala Asn Ser Gly Glu Pro 340 345 350

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Cys Ile Glu 385

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<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:poly-Gly
 flexible linker

<220>

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<223> Gly residues from position 6 to 200 may be present or absent

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